RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/565,371_
Source: _IFWP_
Date Processed by STIC: _1/27/06_

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: _10/565,371_   CRF Edit Date: _1/30/06_
                                    Edited by: _hr_

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

_✓_ Deleted: _✓_ invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

____ Other:

_____
_____
_____

Revised 09/09/2003

IFWP

RAW SEQUENCE LISTING   DATE: 01/30/2006
PATENT APPLICATION: US/10/565,371   TIME: 16:04:02

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\01302006\J565371.raw**

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     5 <110> APPLICANT: TRANSGENE S.A.
     7 <120> TITLE OF INVENTION: Novel multifunctional cytokines
     9 <130> FILE REFERENCE: H2216 PCT S3
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/565,371
C--> 11 <141> CURRENT FILING DATE: 2006-01-23
    11 <150> PRIOR APPLICATION NUMBER: PCT/EP2004/008114
    12 <151> PRIOR FILING DATE: 2004-07-20
    14 <150> PRIOR APPLICATION NUMBER: EP 03 36 0086.7
    15 <151> PRIOR FILING DATE: 2003-07-21
    17 <150> PRIOR APPLICATION NUMBER: US 60/539,320
    18 <151> PRIOR FILING DATE: 2004-01-28
    20 <160> NUMBER OF SEQ ID NOS: 59
    22 <170> SOFTWARE: PatentIn version 3.1
    26 <210> SEQ ID NO: 1
    28 <211> LENGTH: 345
    30 <212> TYPE: PRT
    32 <213> ORGANISM: artificial sequence
    34 <220> FEATURE:
    35 <221> NAME/KEY: source
    36 <223> OTHER INFORMATION: /note= "Description of artificial sequence: fusion human
    37       IL-7/linker/human IL-2"
    41 <400> SEQUENCE: 1
    43 Met Phe His Val Ser Phe Arg Tyr Ile Phe Gly Leu Pro Pro Leu Ile
    44 1               5                   10                  15
    47 Leu Val Leu Leu Pro Val Ala Ser Ser Asp Cys Asp Ile Glu Gly Lys
    48             20                  25                  30
    51 Asp Gly Lys Gln Tyr Glu Ser Val Leu Met Val Ser Ile Asp Gln Leu
    52         35                  40                  45
    55 Leu Asp Ser Met Lys Glu Ile Gly Ser Asn Cys Leu Asn Asn Glu Phe
    56     50                  55                  60
    59 Asn Phe Phe Lys Arg His Ile Cys Asp Ala Asn Lys Glu Gly Met Phe
    60 65                  70                  75                  80
    64 Leu Phe Arg Ala Ala Arg Lys Leu Arg Gln Phe Leu Lys Met Asn Ser
    65             85                  90                  95
    68 Thr Gly Asp Phe Asp Leu His Leu Leu Lys Val Ser Glu Gly Thr Thr
    69             100                 105                 110
    72 Ile Leu Leu Asn Cys Thr Gly Gln Val Lys Gly Arg Lys Pro Ala Ala
    73             115                 120                 125
    76 Leu Gly Glu Ala Gln Pro Thr Lys Ser Leu Glu Glu Asn Lys Ser Leu
    77         130                 135                 140
    80 Lys Glu Gln Lys Lys Leu Asn Asp Leu Cys Phe Leu Lys Arg Leu Leu
    81 145                 150                 155                 160
    84 Gln Glu Ile Lys Thr Cys Trp Asn Lys Ile Leu Met Gly Thr Lys Glu
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RAW SEQUENCE LISTING   DATE: 01/30/2006
PATENT APPLICATION: US/10/565,371   TIME: 16:04:02

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\01302006\J565371.raw**

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85                  165                 170                 175
88 His Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
89              180                 185                 190
92 Met Tyr Arg Met Gln Leu Leu Ser Cys Ile Ala Leu Ser Leu Ala Leu
93          195                 200                 205
96 Val Thr Asn Ser Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln Leu
97      210                 215                 220
100 Gln Leu Glu His Leu Leu Leu Asp Leu Gln Met Ile Leu Asn Gly Ile
101 225                 230                 235                 240
104 Asn Asn Tyr Lys Asn Pro Lys Leu Thr Arg Met Leu Thr Phe Lys Phe
105                 245                 250                 255
108 Tyr Met Pro Lys Lys Ala Thr Glu Leu Lys His Leu Gln Cys Leu Glu
109             260                 265                 270
112 Glu Glu Leu Lys Pro Leu Glu Glu Val Leu Asn Leu Ala Gln Ser Lys
113             275                 280                 285
116 Asn Phe His Leu Arg Pro Arg Asp Leu Ile Ser Asn Ile Asn Val Ile
117     290                 295                 300
120 Val Leu Glu Leu Lys Gly Ser Glu Thr Thr Phe Met Cys Glu Tyr Ala
121 305                 310                 315                 320
124 Asp Glu Thr Ala Thr Ile Val Glu Phe Leu Asn Arg Trp Ile Thr Phe
125                 325                 330                 335
128 Cys Gln Ser Ile Ile Ser Thr Leu Thr
129             340                 345
132 <210> SEQ ID NO: 2
134 <211> LENGTH: 333
136 <212> TYPE: PRT
138 <213> ORGANISM: artificial sequence
140 <220> FEATURE:
141 <221> NAME/KEY: source
142 <223> OTHER INFORMATION: /note= "Description of artificial sequence: fusion murine
143       IL7/linker/murine IL-2"
145 <400> SEQUENCE: 2
147 Met Phe His Val Ser Phe Arg Tyr Ile Phe Gly Ile Pro Pro Leu Ile
148 1               5                   10                  15
151 Leu Val Leu Leu Pro Val Thr Ser Ser Glu Cys His Ile Lys Asp Lys
152             20                  25                  30
155 Glu Gly Lys Ala Tyr Glu Ser Val Leu Met Ile Ser Ile Asp Glu Leu
156         35                  40                  45
159 Asp Lys Met Thr Gly Thr Asp Ser Asn Cys Pro Asn Asn Glu Pro Asn
160     50                  55                  60
163 Phe Phe Arg Lys His Val Cys Asp Asp Thr Lys Glu Ala Ala Phe Leu
164 65                  70                  75                  80
167 Asn Arg Ala Ala Arg Lys Leu Lys Gln Phe Leu Lys Met Asn Ile Ser
168             85                  90                  95
171 Glu Glu Phe Asn Val His Leu Leu Thr Val Ser Gln Gly Thr Gln Thr
172             100                 105                 110
175 Leu Val Asn Cys Thr Ser Lys Glu Glu Lys Asn Val Lys Glu Gln Lys
176     115                 120                 125
179 Lys Asn Asp Ala Cys Phe Leu Lys Arg Leu Leu Arg Glu Ile Lys Thr
```

```
180      130                    135                 140
183 Cys Trp Asn Lys Ile Leu Lys Gly Ser Ile Gly Gly Gly Gly Ser Gly
184 145                    150                 155                 160
187 Gly Gly Gly Ser Met Tyr Ser Met Gln Leu Ala Ser Cys Val Thr Leu
188                    165                 170                 175
191 Thr Leu Val Leu Leu Val Asn Ser Ala Pro Thr Ser Ser Ser Thr Ser
192                180                185                190
195 Ser Ser Thr Ala Glu Ala Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln
196          195                200                205
199 Gln Gln His Leu Glu Gln Leu Leu Met Asp Leu Gln Glu Leu Leu Ser
200      210                215                220
203 Arg Met Glu Asn Tyr Arg Asn Leu Lys Leu Pro Arg Met Leu Thr Phe
204 225                    230                235                240
207 Lys Phe Tyr Leu Pro Lys Gln Ala Thr Glu Leu Lys Asp Leu Gln Cys
208                245                250                255
211 Leu Glu Asp Glu Leu Gly Pro Leu Arg His Val Leu Asp Leu Thr Gln
212          260                265                270
215 Ser Lys Ser Phe Gln Leu Glu Asp Ala Glu Asn Phe Ile Ser Asn Ile
216      275                280                285
219 Arg Val Thr Val Val Lys Leu Lys Gly Ser Asp Asn Thr Phe Glu Cys
220      290                295                300
223 Gln Phe Asp Asp Glu Ser Ala Thr Val Val Asp Phe Leu Arg Arg Trp
224 305                    310                315                320
227 Ile Ala Phe Cys Gln Ser Ile Ile Ser Thr Ser Pro Gln
228                325                330
231 <210> SEQ ID NO: 3
233 <211> LENGTH: 330
235 <212> TYPE: PRT
237 <213> ORGANISM: artificial sequence
239 <220> FEATURE:
240 <221> NAME/KEY: source
241 <223> OTHER INFORMATION: /note= "Description of artificial sequence: fusion human
242         IL-2/linker/human IL-15"
246 <400> SEQUENCE: 3
248 Met Tyr Arg Met Gln Leu Leu Ser Cys Ile Ala Leu Ser Leu Ala Leu
249 1              5                  10                 15
253 Val Thr Asn Ser Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln Leu
254                20                25                 30
257 Gln Leu Glu His Leu Leu Leu Asp Leu Gln Met Ile Leu Asn Gly Ile
258          35                40                 45
261 Asn Asn Tyr Lys Asn Pro Lys Leu Thr Arg Met Leu Thr Phe Lys Phe
262      50                55                 60
265 Tyr Met Pro Lys Lys Ala Thr Glu Leu Lys His Leu Gln Cys Leu Glu
266 65                70                .    75                 80
269 Glu Glu Leu Lys Pro Leu Glu Glu Val Leu Asn Leu Ala Gln Ser Lys
270                85                90                 95
273 Asn Phe His Leu Arg Pro Arg Asp Leu Ile Ser Asn Ile Asn Val Ile
274                100               105               110
277 Val Leu Glu Leu Lys Gly Ser Glu Thr Thr Phe Met Cys Glu Tyr Ala
```

```
278           115                   120                   125
281 Asp Glu Thr Ala Thr Ile Val Glu Phe Leu Asn Arg Trp Ile Thr Phe
282       130                   135                   140
285 Cys Gln Ser Ile Ile Ser Thr Leu Thr Gly Gly Gly Ser Gly Gly
286 145                   150                   155                   160
289 Gly Gly Ser Gly Gly Gly Gly Ser Met Arg Ile Ser Lys Pro His Leu
290                   165                   170                   175
293 Arg Ser Ile Ser Ile Gln Cys Tyr Leu Cys Leu Leu Leu Asn Ser His
294             180                   185                   190
297 Phe Leu Thr Glu Ala Gly Ile His Val Phe Ile Leu Gly Cys Phe Ser
298             195                   200                   205
301 Ala Gly Leu Pro Lys Thr Glu Ala Asn Trp Val Asn Val Ile Ser Asp
302       210                   215                   220
305 Leu Lys Lys Ile Glu Asp Leu Ile Gln Ser Met His Ile Asp Ala Thr
306 225                   230                   235                   240
309 Leu Tyr Thr Glu Ser Asp Val His Pro Ser Cys Lys Val Thr Ala Met
310                   245                   250                   255
313 Lys Cys Phe Leu Leu Glu Leu Gln Val Ile Ser Leu Glu Ser Gly Asp
314             260                   265                   270
317 Ala Ser Ile His Asp Thr Val Glu Asn Leu Ile Ile Leu Ala Asn Asn
318       275                   280                   285
321 Ser Leu Ser Ser Asn Gly Asn Val Thr Glu Ser Gly Cys Lys Glu Cys
322       290                   295                   300
325 Glu Glu Leu Glu Glu Lys Asn Ile Lys Glu Phe Leu Gln Ser Phe Val
326 305                   310                   315                   320
329 His Ile Val Gln Met Phe Ile Asn Thr Ser
330                   325                   330
333 <210> SEQ ID NO: 4
335 <211> LENGTH: 330
337 <212> TYPE: PRT
339 <213> ORGANISM: artificial sequence
341 <220> FEATURE:
342 <221> NAME/KEY: source
343 <223> OTHER INFORMATION: /note= "Description of artificial sequence: fusion human
344       IL-15/linker/human IL-2"
348 <400> SEQUENCE: 4
350 Met Arg Ile Ser Lys Pro His Leu Arg Ser Ile Ser Ile Gln Cys Tyr
351 1               5                   10                   15
354 Leu Cys Leu Leu Leu Asn Ser His Phe Leu Thr Glu Ala Gly Ile His
355             20                   25                   30
358 Val Phe Ile Leu Gly Cys Phe Ser Ala Gly Leu Pro Lys Thr Glu Ala
359             35                   40                   45
362 Asn Trp Val Asn Val Ile Ser Asp Leu Lys Lys Ile Glu Asp Leu Ile
363       50                   55                   60
366 Gln Ser Met His Ile Asp Ala Thr Leu Tyr Thr Glu Ser Asp Val His
367 65                   70                   75                   80
370 Pro Ser Cys Lys Val Thr Ala Met Lys Cys Phe Leu Leu Glu Leu Gln
371                   85                   90                   95
374 Val Ile Ser Leu Glu Ser Gly Asp Ala Ser Ile His Asp Thr Val Glu
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RAW SEQUENCE LISTING   DATE: 01/30/2006
PATENT APPLICATION: US/10/565,371   TIME: 16:04:02

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\01302006\J565371.raw

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375              100                105                110
379 Asn Leu Ile Ile Leu Ala Asn Asn Ser Leu Ser Ser Asn Gly Asn Val
380          115                120                125
383 Thr Glu Ser Gly Cys Lys Glu Cys Glu Glu Leu Glu Glu Lys Asn Ile
384      130                135                140
387 Lys Glu Phe Leu Gln Ser Phe Val His Ile Val Gln Met Phe Ile Asn
388 145                150                155                160
391 Thr Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
392              165                170                175
395 Ser Met Tyr Arg Met Gln Leu Leu Ser Cys Ile Ala Leu Ser Leu Ala
396          180                185                190
399 Leu Val Thr Asn Ser Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln
400          195                200                205
404 Leu Gln Leu Glu His Leu Leu Leu Asp Leu Gln Met Ile Leu Asn Gly
405      210                215                220
408 Ile Asn Asn Tyr Lys Asn Pro Lys Leu Thr Arg Met Leu Thr Phe Lys
409 225                230                235                240
412 Phe Tyr Met Pro Lys Lys Ala Thr Glu Leu Lys His Leu Gln Cys Leu
413              245                250                255
416 Glu Glu Glu Leu Lys Pro Leu Glu Glu Val Leu Asn Leu Ala Gln Ser
417          260                265                270
420 Lys Asn Phe His Leu Arg Pro Arg Asp Leu Ile Ser Asn Ile Asn Val
421          275                280                285
424 Ile Val Leu Glu Leu Lys Gly Ser Glu Thr Thr Phe Met Cys Glu Tyr
425      290                295                300
428 Ala Asp Glu Thr Ala Thr Ile Val Glu Phe Leu Asn Arg Trp Ile Thr
429 305                310                315                320
432 Phe Cys Gln Ser Ile Ile Ser Thr Leu Thr
433              325                330
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436 <210> SEQ ID NO: 5
438 <211> LENGTH: 350
440 <212> TYPE: PRT
442 <213> ORGANISM: artificial sequence
444 <220> FEATURE:
445 <221> NAME/KEY: source
446 <223> OTHER INFORMATION: /note= "Description of artificial sequence: fusion signal IL-2/
447   human IL-15/linker/human IL-2"
451 <400> SEQUENCE: 5

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453 Met Tyr Arg Met Gln Leu Leu Ser Cys Ile Ala Leu Ser Leu Ala Leu
454 1              5                10                15
457 Val Thr Asn Ser Met Arg Ile Ser Lys Pro His Leu Arg Ser Ile Ser
458              20                25                30
461 Ile Gln Cys Tyr Leu Cys Leu Leu Leu Asn Ser His Phe Leu Thr Glu
462          35                40                45
465 Ala Gly Ile His Val Phe Ile Leu Gly Cys Phe Ser Ala Gly Leu Pro
466      50                55                60
469 Lys Thr Glu Ala Asn Trp Val Asn Val Ile Ser Asp Leu Lys Lys Ile
470 65                70                75                80
473 Glu Asp Leu Ile Gln Ser Met His Ile Asp Ala Thr Leu Tyr Thr Glu
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VERIFICATION SUMMARY   DATE: 01/30/2006
PATENT APPLICATION: **US/10/565,371**   TIME: 16:04:03

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\01302006\J565371.raw**

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

**Raw Sequence Listing before editing,
for reference only**

IFWP

RAW SEQUENCE LISTING   DATE: 01/27/2006
PATENT APPLICATION: US/10/565,371   TIME: 14:34:46

Input Set : A:\H2216PCT_correctedsequence.txt
Output Set: N:\CRF4\01272006\J565371.raw

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  5 <110> APPLICANT: TRANSGENE S.A.
  7 <120> TITLE OF INVENTION: Novel multifunctional cytokines
  9 <130> FILE REFERENCE: H2216 PCT S3
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/565,371
C--> 12 <141> CURRENT FILING DATE: 2006-01-23
  14 <150> PRIOR APPLICATION NUMBER: EP 03 36 0086.7
  15 <151> PRIOR FILING DATE: 2003-07-21
  17 <150> PRIOR APPLICATION NUMBER: US 60/539,320
  18 <151> PRIOR FILING DATE: 2004-01-28
  20 <160> NUMBER OF SEQ ID NOS: 59
  22 <170> SOFTWARE: PatentIn version 3.1
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ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed

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2761 <210> SEQ ID NO: 59
2763 <211> LENGTH: 49
2765 <212> TYPE: DNA
2767 <213> ORGANISM: artificial sequence
2769 <220> FEATURE:
2770 <221> NAME/KEY: source
2771 <223> OTHER INFORMATION: /note= "Description of artificial sequence: 3' linker
primer
2772        for generating the mIL15/IL2 fusion"
2776 <400> SEQUENCE: 59
2777 agagccacct ccgcctgaac cgcctccacc cttgtcatcg tcgtccttg          49
E--> 2780 (1)
E--> 2783 (45)
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VERIFICATION SUMMARY   DATE: 01/27/2006
PATENT APPLICATION: **US/10/565,371**   TIME: 14:34:47

Input Set : **A:\H2216PCT_correctedsequence.txt**
Output Set: **N:\CRF4\01272006\J565371.raw**

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2780 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=59